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(72) Inventors; and
(75) Inventors/Applicants (for US only): **WEBSTER, Mitch** [ZA/NZ]; 161 Kennedy Road, Albany R D 2, AUCKLAND (NZ). **HORLACHER, Oliver** [NZ/NZ]; 7/10 Tagalad Road, Mission Bay, AUCKLAND (NZ).

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(71) Applicant (for all designated States except US): **CARSHA COMPANY LIMITED** [NZ/NZ]; 161 Kennedy Road, Albany R D 2, AUCKLAND (NZ).

(74) Agents: **ADAMS, Matthew, D** et al.; A J Park, 6th Floor Huddart Parker Building, PO Box 949, Wellington, 6015 (NZ).

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(54) Title: METHODS FOR PROCESSING GENOMIC INFORMATION AND USES THEREOF

A
Randomise the list

B

Annotation

Name	Chromosome	Copies	Start	Stop
CYP2D6	3	1	56	1065
ACO1	5	1	7865	8763
HIA1	12	1	12	2000
ABCA6	x	1	5748	6003

Number list

Annotation

Name	Chromosome	Copies	Start	Stop
1 CYP2D6	3	1	56	1065
2 ACO1	5	1	7865	8763
3 HIA1	12	1	12	2000
4 ABCA6	x	1	5748	6003

Separate the datasets of gene names and gene annotation information.

Gene names

1 CYP2D6
2 ACO1
3 HIA1
4 ABCA6

Gene data

1 3, 1, 56, 1065
2 5, 1, 7865, 8763
3 12, 1, 12, 2000
4 x, 1, 5748, 6003

(57) Abstract: Methods for processing and storing genomic information in a secure manner are described. In particular, methods for processing, splitting and storing genomic information or portions thereof are disclosed. An individual's genomic information is digitized and a splitting algorithm applied to fragment and randomise the digitized genomic information into at least two separate datasets. Access to at least one dataset is retained by the individual and the second dataset is stored on a central server as a secure database record. Each dataset in isolation presents uninformative data and it is only when all datasets are combined that the data is capable of being presented into a useable and informative format.

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